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## COMMUNICATION

## Gelation by supramolecular dimerization of mono(urea)s

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**Mono-ureido salicylic acid derivatives form hydrogen bonded dimers that mimic the gelation behaviour of covalently linked bis(urea)s. The 5-ureido salicylic acid isomers are much more effective gelators because of the lowered conjugation and less planar geometry, resulting in stronger urea...urea  $\alpha$ -tape hydrogen bonding interactions.**

### Introduction

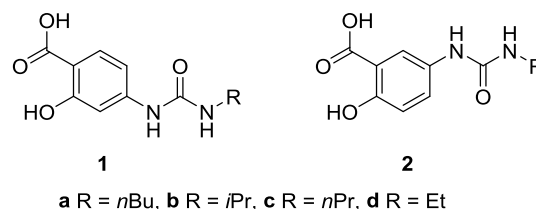
Gel phase soft materials based on supramolecular interactions between low molecular weight gelators (LMWG) have become extremely popular in the past decade. These versatile media have applications<sup>1,2</sup> in organocatalysis,<sup>3,4</sup> wound healing,<sup>5</sup> drug delivery,<sup>6,7</sup> as templates and supports for nanomaterials,<sup>8-10</sup> and in pharmaceutical polymorph screening and control.<sup>11-15</sup> LMWG represent a very broad class of compounds which have in common the ability to form high aspect ratio fibres by rapid unidirectional growth of a solid, sample-spanning network. Gels are formed in a wide range of solvents including water (hydrogels) and both polar and non-polar organic liquids (organogels). The reversibility of the supramolecular interactions between the small molecule gelators means that gels derived from LMWG are usually thermally reversible and their materials properties, particularly their rheological characteristics, can be manipulated by light, changes in pH or by chemical triggers such as metal cations and hydrogen bond acceptor anions.<sup>16-21</sup> Common LMWG are based on chemical functionalities capable of forming one dimensional assemblies including nucleobases, amides and ureas.<sup>21-23</sup> Most LMWG incorporate multiple functional groups that associate in a directional way in order to increase the strength of the fibres by multivalent supramolecular interactions. In urea-based systems most gelators (with some exceptions<sup>24-27</sup>) contain at least two urea groups.<sup>25, 28-35</sup> We now report a novel approach to effective urea-based gelators in which two urea functional groups are linked by a supramolecular hydrogen bonded motif instead of by a covalent bond. The interactions involved in the assembly of the mono(urea) LMWG into a dimer and the subsequent assembly of the hydrogen bonded dimers into a fibre are orthogonal to one another, and the different supramolecular

homosynthons form with high fidelity, allowing effective gelation. The approach is related to the hexagonal isophthalic acid derived assemblies reported by Potluri and Hamilton.<sup>24</sup>

### Results and Discussion

#### Gelator synthesis

A series of mono(urea)s based on 4- and 5-aminosalicylic acid were synthesised from reaction of the respective aminosalicylic acid with the appropriate isocyanate to give compounds **1a-d** and **2a-d** in good yield. A search of the CSD shows that the salicylic acid functionality is almost invariably planar because of strong intramolecular OH...O hydrogen bonding and tends to form hydrogen bonded carboxylic acid dimers *via* an  $R_2^2(8)$  hydrogen bonded ring motif,<sup>36</sup> as in the structure of 5-aminosalicylic acid itself and 17 out of 24 related derivatives.<sup>37</sup> Based on the CSD evidence the supramolecular synthons<sup>38</sup> formed by the salicylic acid and urea functional groups might be expected to be orthogonal to one another and hence self-sorting in their assembly behaviour.



#### Gelation

Gelation tests on compounds of type **1** and **2** were carried out at 1 mg mL<sup>-1</sup> in a wide range of solvents. Samples were sonicated for 30 seconds, then heated gently to dissolve the solid, and subsequently allowed to cool to room temperature. Most gels formed within a few minutes while for some samples crystallisation occurred over an approximately 24 hour period. Full results are tabulated in the supplementary information, Table S1. The 5-ASA derivatives of type **2** proved to be surprisingly efficient gelators despite the presence of only a single urea group. The unbranched chain members of the series **2a, c** and **d** form gels to some degree in 1,2,4-

trichlorobenzene, 1,2-dibromoethane, 1,2-dichlorobenzene, 1,3-dichlorobenzene and nitrobenzene (see supplementary Figure S1). The gels form within a few minutes of cooling and are all opaque suggesting a relatively large particle size. Gels formed in 1,2,4-trichlorobenzene collapsed after 30 min, while 1,2-dibromoethane and 1,3-dichlorobenzene gave somewhat stronger gels, although these also collapsed after a few hours. The strongest and most optically transparent gels formed in 1,2-dichlorobenzene and nitrobenzene, which proved to be stable at least over a period of weeks. In comparison with the unbranched gelators, the isopropyl derivative **2b** formed only partial gels in just two solvents and these materials collapsed immediately. It is possible that the increased steric bulk of the isopropyl group hinders urea self-assembly and hence disfavours gel formation.

The 4-ASA derivatives of type **1** proved to be more soluble than their 5-ASA counterparts and did not exhibit significant gelation behaviour. In only one case was a gel-like material obtained. Slow cooling of **1a** in nitromethane gave an opaque, white self-supporting solid with a critical gelation concentration of 1% w/v. SEM analysis of the resulting xerogel showed it to comprise a mixture of fibrous and crystalline material (see supplementary information, Figures S1 and S2).

Representative gels were dried in a vacuum desiccator for 24 hours to give xerogels and then coated with either chromium or platinum and imaged by SEM. The microscope images showed that the xerogels adopt the expected fibrous morphology. Fibre widths and bunching vary from sample to sample but are consistent with other semi-transparent gels.<sup>39</sup> A representative image of the xerogel of *n*-butyl compound **2a** obtained from nitrobenzene is shown in Figure 1a and contrasts with the xerogel of ethyl analogue **2d**, which exhibits a mixture of shorter, blockier and more well defined crystalline tapes and long, thin and more curved fibres (Figure 1b). The shorter ethyl substituent may result in more facile recrystallization of some of the sample during the drying process. SEM images of nitromethane xerogels of **1a**, the only 4-ASA derived gel, show a significant difference from the 5-ASA versions, with wide crystalline aggregates of fibres forming sheets 1  $\mu$ m across (supplementary information Figure S4). All the fibres are distinctly larger than those of **1a** and **1d** in the same solvent. This may account for the more crystalline, opaque appearance of the gel.

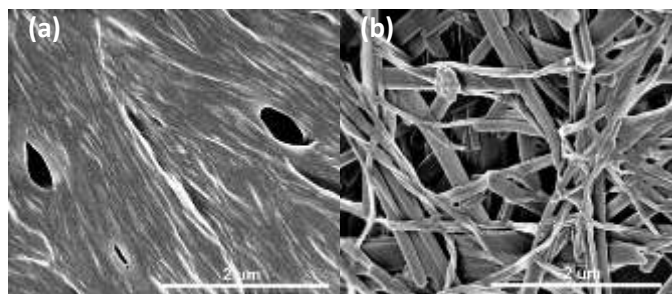


Figure 1. SEM images of (a) xerogel of **2a** obtained from nitrobenzene and (b) xerogel of **2d** obtained from nitromethane.

Gels of **2a**, **2c** and **2d** were characterised by oscillatory stress sweep rheometry to probe their viscoelastic properties in several solvents. The solid-like character of the materials was manifest in the fact that the  $G'$  (storage modulus) remains well over an order of magnitude higher than the viscous modulus  $G''$  up to the yield stress of the gel. The relatively weak nature of the gels of compound **2a** in 1,2-dichlorobenzene is apparent in the fairly low value of  $G'$  of ca. 1000

Pa at low stress, and the low yield stress of around 20 Pa, for example (Figure 2a). This gel exhibits weak strain overshoot characterised by a rise on  $G''$  with increasing stress. This viscoelastic behaviour is indicative of close-packed weakly bonded particles forming a structure that resists deformation just prior to breaking of interactions and flow occurring.<sup>40, 41</sup> Elastic moduli were found to be highly solvent dependent and to increase with increasing chain length. Hence, 1,2-dibromoethane gels of the butyl compound **2a** proved quite robust with  $G'$  reaching ca. 10 kPa (Figure 2b, see supplementary information Figure S5).

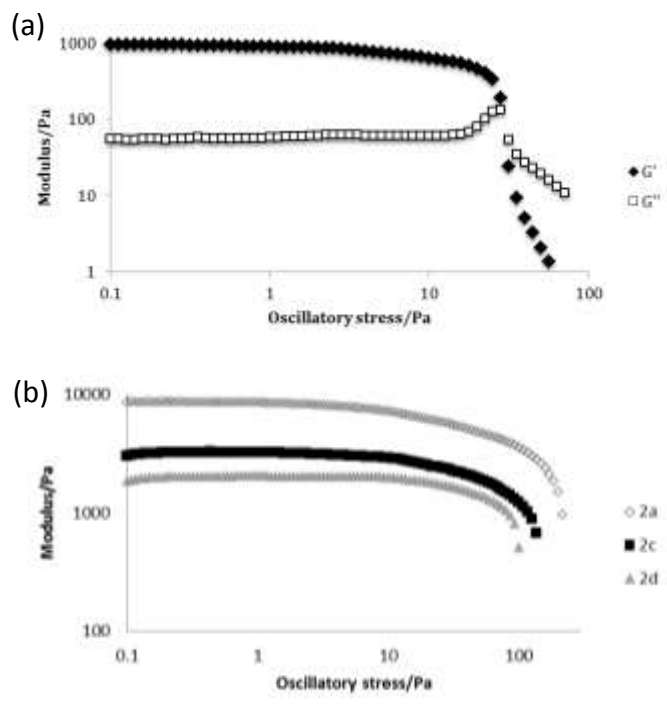


Figure 2. (a) Stress sweep rheometry for a 1% w/v gel of **2a** in 1,2-dichlorobenzene. (b) comparison of the evolution of  $G'$  in 1,2-dibromoethane as a function of chain length.

Single crystals suitable for X-ray crystallography were obtained for compounds **1a**, **1c** and **1d** by recrystallization from ethyl acetate, acetone and 2-butanone, respectively. The sample of **1d** proved to be of very low quality and is not reported herein but a qualitative structure solution confirmed the packing is similar to the other two compounds of type **1**. The structures of **1a** and **1c** are of high precision. The structures of the 4-ASA derivatives of type **1** all exhibit the common head-to-head carboxylic acid dimer motif,<sup>42</sup> with approximate co-planarity of the carboxylic acid group and aryl ring enforced by an intramolecular OH...O interaction from the phenolic OH group to the acid carbonyl functionality, Figure 3. The high acidity of the salicylic acid moiety results in short O...O distances across the  $R_2^2(8)$  hydrogen bonded ring of 2.61 – 2.65 Å. The urea moieties are not precisely co-planar with the salicylic acid ring although there is a relatively long intramolecular ArCH...O=C hydrogen bond, as is the case with electron withdrawing diarylureas.<sup>43, 44</sup> In **1a** the urea groups form a twisted  $R_1^2(6)$  urea  $\alpha$ -tape hydrogen bonded chain that is independent of the acid dimer hydrogen bonding. This urea motif, particularly in an antiparallel arrangement in covalently linked bis(urea)s, has been linked to the ubiquitous gel-forming ability of these compounds.<sup>28, 39, 45</sup> However, in **1a**, each hydrogen bonded dimer links into two different twisted  $\alpha$ -tapes to give an overall 2D hydrogen bonded network based on

two crystallographically independent molecules. In **1c**, in contrast, the two urea  $\alpha$ -tapes formed by each hydrogen bonded dimer are approximately coplanar and align antiparallel to give a 1D fibre, mimicking the arrangement found in covalently bonded bis(ureas).

The 5-ASA derivative **2a** was also characterised by single crystal X-ray crystallography using a sample obtained by recrystallization from methanol, Figure 3c. As with the compounds of type **1**, this compound also exhibits a head-to-head carboxylic acid dimer arrangement and, as with **1c**, each dimer stacks with its neighbours to give an antiparallel pair of urea  $\alpha$ -tape interactions running throughout the crystal. Notably the urea  $\text{NH}\cdots\text{O}$  distances of 2.80 – 2.82 Å are at the lower end of the typical range found for this interaction<sup>44</sup> and are considerably shorter than the in **1a** and **1c** (the interactions in **1a** are unsymmetrical at 2.81/3.07 and 2.88/2.98 Å, while those in **1c** are symmetrical at 2.89 – 2.91 Å). This short urea $\cdots$ urea distance is correlated with the much more significant out-of-plane orientation of the urea group relative to the salicylic acid, which breaks the intramolecular  $\text{CH}\cdots\text{O}$  interaction.

It is plausible that gelation by compounds of type **1** and particularly type **2** arises from urea  $\alpha$ -tape formation, and the fact that this interaction is much shorter in **2a** is consistent with the much more effective gelation behaviour of this compound compared to compounds of type **1**. While mono(urea)s interact with one another through a single bifurcated acceptor  $R_2^1(6)$  synthon, covalently bound bis(urea)s enjoy a multivalent<sup>46</sup> stabilization by virtue of the double  $R_2^1(6)$  interaction formed by each molecule. In the present case we postulate that the strong non-covalent dimerization arising from the salicylic acid groups in compounds of type **1** and **2** allows them to behave as *pseudo*-bis(urea)s with the urea groups in hydrogen bonded pairs held in a centrosymmetric and hence antiparallel arrangement by the centrosymmetric  $R_2^2(8)$  carboxylic acid dimer arrangement. Evidence for the persistence of the dimeric structure in the gas phase comes from the  $\text{ES}^+$  mass spectrum of **2a** which, in addition to the molecular ion peak at  $m/z$  252.8, exhibits a distinct peak at 505.1 of approximately half the intensity of the molecular ion, assigned to the dimer. Similar gas phase dimers are readily apparent in the  $\text{ES}^+$ -MS spectra of all compounds of type **1** and **2**. In solution the  $^1\text{H}$  NMR spectrum of **2a** in  $\text{DMSO}-d_6$  solution did not exhibit any change in the chemical shift of the carboxylic acid OH resonance as a function of temperature in the range 20 – 100 °C, nor in the concentration range 1 – 8% w/v. In contrast, a modest chemical shift change was observed for the NH resonances, suggesting that while aggregation by urea-urea interaction is in equilibrium under the conditions used in the NMR experiment, the carboxylic acid groups remain fully dimerised under the conditions studied even in a competitive solvent such as DMSO. It is also possible that the lack of chemical shift change as a function of concentration and temperature indicates no interaction at all and the molecule is monomeric; however, this seems unlikely given the strong hydrogen bonding in the salicylic acid group, the  $\text{ES}^+$ -MS evidence and the contrasting behaviour of the generally more weakly hydrogen bonded urea group.<sup>47</sup> Definitive proof for the retention of the dimeric structure in solution was obtained from IR spectroscopy. The solid-state ATR spectra of all compounds of type **1** and **2** closely resembled the spectra in the non-competitive solvent 1,2-dibromoethane with a single low frequency asymmetric carbonyl stretching band at 1660  $\text{cm}^{-1}$  for **2a**, for example, assigned to the carboxylic acid dimer observed in the X-ray crystal structures. In the more competitive DMSO solution, however, two carboxylic acid carbonyl bands appeared, at 1703 and 1655  $\text{cm}^{-1}$  assigned to monomeric and dimeric carboxylic acids respectively. Other consistent changes were observed throughout the spectra. Hence in

the competitive solvent the compound exists as a mixture of monomer and dimer while in non-polar media it is fully dimerised.

Evidence for the relevance of the single crystal X-ray structural data to the gels formed by compounds of type **2** was sought from X-ray powder diffraction.<sup>48-50</sup> However the XRPD patterns of the xerogels of **2a**, **2c** and **2d** from nitromethane and nitrobenzene all showed the materials to be amorphous. Given the simplicity of the compounds and the consistent structural results, however, it remains a plausible working hypothesis that the local structure of the gel fibres is closely related to the interactions found in the single crystals.

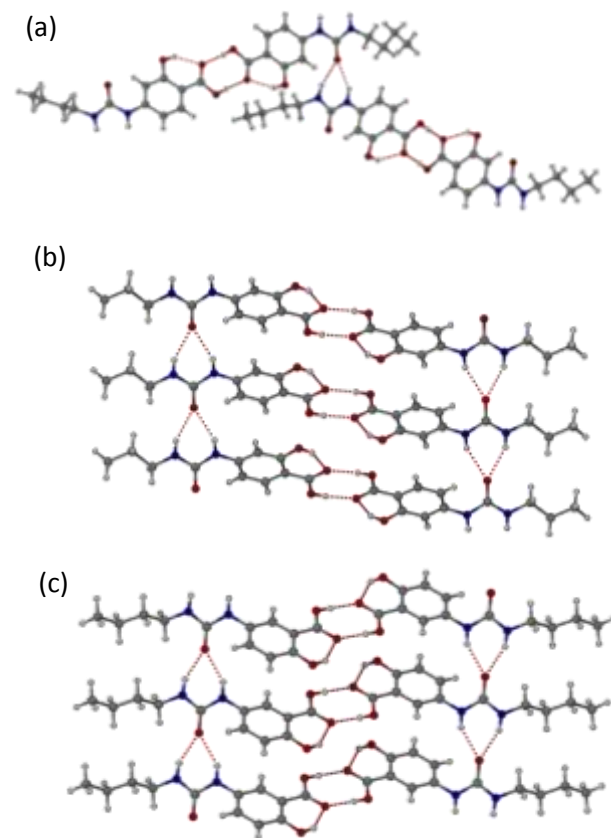


Figure 3. (a) Twisted 2D urea hydrogen bonding in **1a**; (b) parallel packing in **1c** with relatively long  $\text{NH}\cdots\text{O}$  interactions; (c) short  $\text{NH}\cdots\text{O}$  interactions in the gelator **2a**. Selected hydrogen bonding distances (Å) for compound **1a**: Acid dimer  $\text{O}\cdots\text{O}$  2.606(2), 2.634(2); urea tape  $\text{N}\cdots\text{O}$  2.986(3), 2.880(2), 2.814(2) and 3.072(2), compound **1c**,  $\text{O}\cdots\text{O}$  2.892(3),  $\text{N}\cdots\text{O}$  2.911(3), 2.892(3); compound **2a**,  $\text{O}\cdots\text{O}$  2.629(6);  $\text{N}\cdots\text{O}$  2.797(7), 2.816(7).

Given that both the 4-ASA derivatives of type **1** and 5-ASA analogues of type **2** share a common dimeric structure, it is interesting to see that compounds of type **2** exhibit shorter urea tape hydrogen bonding and are also significantly more effective gelators of the solvents studied than compounds of type **1**. The out-of-plane twist of the urea group compared to the aromatic ring plane in **2a** is exemplified by a torsion angle of 50.9°, significantly larger than those of the 4-ASA derivatives which are between 17.3 – 38.0°. Gas-phase DFT calculations suggest that this is not solely a crystal packing effect, with the rotation energy barrier for the 4-ASA derivatives being around 29  $\text{kJ mol}^{-1}$ , nearly double the analogous barrier for the 5-ASA analogues (supplementary information Figure

S6). This more restricted rotation is likely to be due to the greater conjugation of the urea nitrogen atom in the 4-ASA derivative (Figure S8) as a result of the para-electron withdrawing effect of the carboxylic acid functionality.<sup>51</sup> As a result the hydrogen bond acceptor ability of the urea carbonyl group is reduced for steric reasons and hence its ability to express the gel-forming urea  $\alpha$ -tape motif is limited.<sup>45, 52</sup> Overall, the short NH...O interactions in X-ray structure of **2a** indicate that the urea  $\alpha$ -tape is significantly stronger than those displayed by the 4-ASA analogues of type 1. Interestingly, this electron withdrawing effect is evident in the structure of 4-ASA itself which is neutral, while 5-ASA is a zwitterion because of the higher amine basicity.<sup>37</sup>

## Conclusions

In conclusion this work has shown that a robust *supramolecular* linkage can replace a covalently bonded spacer in bis(urea)s, allowing mono(urea)s to act as effective gelators by means of supramolecular multivalency. In addition to the formation of multiple hydrogen bonded chain interactions, gelation is favoured by non-coplanarity of the urea group and the aryl substituent, maximising the exposure and hence hydrogen bond basicity of the urea carbonyl acceptor. The presence of sterically bulky branched substituents as in **2b** also apparently reduces gelation efficiency, again likely as a result of reduced steric accessibility of the urea carbonyl group and hence reduced hydrogen bonded chain growth rate.

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## Notes and references

a Department of Chemistry, South Road, University of Durham, DH1 3LE (UK). E-mail: jon.steed@durham.ac.uk; Fax: +44(0)191 384 4737; Tel: +44 (0)191 334 2085. Electronic Supplementary Information (ESI) available: tables of gelation experiments and single crystal structure determination in CIF format CCDC 1419680-1419682. See DOI: 10.1039/c000000x/ The underlying research data for this paper is available in accordance with EPSRC open data policy from <http://dx.doi.org/10.15128/kk91fm27z>.

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## Graphical Abstract

